

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: October 5, 2005, 06:38:44 ; Search time 44 Seconds
(without alignments)

816.050 Million cell updates/sec

Title: US-09-884-260A-7

Perfect score: 2488 MATRSSSSPELPKIPKPGY.....DPLDPAVWFKSLTRATNY 481

Sequence: 1 MATRSSSSPELPKIPKPGY.....DPLDPAVWFKSLTRATNY 481

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Listing filter 45 summaries

Database : Issued Patents AA.*
1: /cgn2_6/prodata/1/1aa/5a_COMB pep.*
2: /cgn2_6/prodata/1/1aa/5b_COMB pep.*
3: /cgn2_6/prodata/1/1aa/6a_COMB pep.*
4: /cgn2_6/prodata/1/1aa/6b_COMB pep.*
5: /cgn2_6/prodata/1/1aa/PCUS_COMB pep.*
6: /cgn2_6/prodata/1/1aa/backfile1 pep.*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed.
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2488	100.0	481	3	US-09-537-357-7
2	2463.5	99.0	487	3	US-09-537-357-15
3	965.5	38.8	483	3	US-08-833-553-2
4	965.5	38.8	483	3	US-09-418-222-2
5	965.5	38.8	483	3	US-09-537-357-36
6	961.5	38.6	483	3	US-08-878-1734-6
7	961.5	38.6	483	3	US-08-878-1734-5
8	961.5	38.6	483	3	US-08-878-1734-25
9	961.5	38.6	483	3	US-09-537-357-25
10	961.5	38.6	483	3	US-09-537-357-12
11	961.5	38.6	483	3	US-09-537-357-12
12	961.5	38.6	483	3	US-09-537-357-12
13	961.5	38.6	483	3	US-09-537-357-12
14	961.5	38.6	483	3	US-09-537-357-12
15	961.5	38.6	483	3	US-09-537-357-12
16	961.5	38.6	483	3	US-09-537-357-12
17	961.5	38.6	483	3	US-09-537-357-12
18	961.5	38.6	483	3	US-09-537-357-12
19	961.5	38.6	483	3	US-09-537-357-12
20	961.5	38.6	483	3	US-09-537-357-12
21	961.5	38.6	483	3	US-09-537-357-12
22	961.5	38.6	483	3	US-09-537-357-12
23	961.5	38.6	483	3	US-09-537-357-12
24	961.5	38.6	483	3	US-09-537-357-12
25	961.5	38.6	483	3	US-09-537-357-12
26	961.5	38.6	483	3	US-09-537-357-12
27	961.5	38.6	483	3	US-09-537-357-12

28	184	7.4	62	3	US-09-537-357-11	Sequence 11, Appl
29	183	7.4	68	3	US-08-833-553-9	Sequence 9, Appl
30	183	7.4	68	3	US-09-418-222-9	Sequence 9, Appl
31	179.5	7.2	496	4	US-10-142-231-90	Sequence 90, Appl
32	179.5	7.2	503	4	US-09-583-447A-2	Sequence 2, Appl
33	175.5	7.1	61	3	US-09-537-357-10	Sequence 10, Appl
34	168.5	6.8	504	4	US-09-583-447A-4	Sequence 4, Appl
35	166.5	6.7	420	4	US-09-583-447A-4	Sequence 4, Appl
36	164.5	6.6	512	4	US-09-583-447A-4	Sequence 4, Appl
37	160	6.4	488	1	US-08-470-757-5399	Sequence 6, Appl
38	160	6.4	488	1	US-08-470-757-5399	Sequence 6, Appl
39	158	6.4	488	1	US-08-470-757-5399	Sequence 6, Appl
40	158	6.4	488	1	US-08-470-757-5399	Sequence 6, Appl
41	158	6.4	488	1	US-08-470-757-5399	Sequence 6, Appl
42	156.5	6.3	500	4	US-10-142-231-61	Sequence 61, Appl
43	156	6.3	32	3	US-09-537-357-30	Sequence 30, Appl
44	156	6.3	507	1	US-08-457-274A-23	Sequence 23, Appl
45	156	6.3	507	5	PCT-US95-05758-23	Sequence 23, Appl

ALIGNMENTS

US-09-537-357-7	Sequence 7, Appl	Application US/09537357
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45	45	45

Db	361	VGGHSSSPKIKKKGIITFGYPPATADPKIFDSEKFGVGRFVGEGBEGLINKVYMSNR	420
Qy	421	ETPEPSPNNCCGCGCNVYIGRIWEPFPLDTFTFVADPLGSAVYKSLFRATDM	480
Db	421	ETPEPSPNNCCGCGCNVYIGRIWEPFPLDTFTFVADPLGSAVYKSLFRATDM	480
Qy	481	V	481
Db	481	V	481

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1  RESULT: 2
2  US-09-537-357-15
3  Patent No. 6271018
4  Title of Invention: Application US/09517357
5  GENERAL INFORMATION:
6  APPLICANT: Alan Brash
7  TITLE OF INVENTION: MUSKIELOX (CUCUMIS MELO) HYDROPEROXIDE
8  FILE REFERENCE: 06037.0002 US/09/537.357
9  CURRENT APPLICATION NUMBER: 09-03-25
10 NUMBER OF SEQ ID NOS: 56
11 SOFTWARE: FastSeq for Windows Version 4.0
12 SEQ ID NO 15
13 LENGTH: 487
14 TYPE: PRT
15 ORGANISM: Cucumis melo
16 NAME: MUSKIELOX
17 LOCATION: (1)
18 OTHER INFORMATION: Xaa = Any Amino Acid
19 NAME/KEY: misc feature
20 LOCATION: (0)...(0)
21 OTHER INFORMATION: Accession No. 6271018 AF081955
22 US-09-537-357-15

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Query Match	99.04;	Score 2463.5;	DB 3;	Length 487;
Best Local Similarity	99.24;	Pred. No. 2.2e-243;		
Matches 478; Conservative	1;	Mismatches 2;	Indels 1;	Gaps 1

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Db 1 MATBSSSSBBLDLEB HGGVBPBPALBP KORVDYFQGHDEP RSNTKNSYTFYRN 50
QY 1 MATBSSSSBBLDLEB HGGVBPBPALBP KORVDYFQGHDEP RSNTKNSYTFYRN 50
Db 1 MATBSSSSBBLDLEB HGGVBPBPALBP KORVDYFQGHDEP RSNTKNSYTFYRN 50
QY 120 ETEHSHVLRFLFSLASBRDDEFPFLFRSSLSSEMPVLRBTKSRKKLADNNSISDSNFD 179
Db 60 MPBPBPSSSSSNVYVLLDLSFPLLPDPAKEXNITDQTMPSLSFPGNIRKTCALDSS 119
QY 61 MPBPBPSSSSSNVYVLLDLSFPLLPDPAKEXNITDQTMPSLSFPGNIRKTCALDSS 120
Db 61 MPBPBPSSSSSNVYVLLDLSFPLLPDPAKEXNITDQTMPSLSFPGNIRKTCALDSS 120
QY 121 ETEHSHVLRFLFSLASBRDDEFPFLFRSSLSSEMPVLRBTKSRKKLADNNSISDSNFD 180
Db 121 ETEHSHVLRFLFSLASBRDDEFPFLFRSSLSSEMPVLRBTKSRKKLADNNSISDSNFD 180
QY 180 YVFRLLSGTPGOSPLAAERQWMDLWYQALPLASISGPGYFVDEYVHTPLPLPFP 239
Db 181 YVFRLLSGTPGOSPLAAERQWMDLWYQALPLASISGPGYFVDEYVHTPLPLPFP 240
QY 240 VFSGGRKYIYAAYSSSSQSLDEAEKQIDIEAQCYNVLAGNAYGKGMVFPPLLKKV 239
Db 241 VFSGGRKYIYAAYSSSSQSLDEAEKQIDIEAQCYNVLAGNAYGKGMVFPPLLKKV 240
QY 300 GTAGEDLHRLAEVRITVNEBSGLTFSALREKSLKSVTEALRIEPPVOTGAKED 359
Db 301 GTAGEDLHRLAEVRITVNEBSGLTFSALREKSLKSVTEALRIEPPVOTGAKED 360
QY 360 IVLOSDBSPKLIKGETFYGVPATPKPIKQSEKFGVPEBEBELKATVYMSSE 419
Db 361 IVLOSDBSPKLIKGETFYGVPATPKPIKQSEKFGVPEBEBELKATVYMSSE 420
QY 420 RETVEYTEAKKCCPGRKLVYIGRIINNVBPFLARDTFTYVADLPAGAVKSLTRATD 479
Db 421 RETVEYTEAKKCCPGRKLVYIGRIINNVBPFLARDTFTYVADLPAGAVKSLTRATD 480

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OY      480 MV 481
DB      481 ML 482

RESULT 3
US-08-833-553-2
; Sequence 2, Application US/08833553C
; Patent No. 6008034
; GENERAL INFORMATION:
; INVENTOR: Alex
; APPLICANT: Leitch Konrad
; APPLICANT: Mubheim, Andreas
; APPLICANT: Silke, Nataasha
; TITLE OF INVENTION: HYDROPEROXIDE LYASES
; FILE REFERENCE: Hydroperoxide Lyases
; CURRENT APPLICATION NUMBER: US/08/833,553C
; CURRENT FILING DATE: 1997-04-07
; NUMBER OF SEQ. ID NOS.: 2
; SOFTWARE: PatentIn Ver. 2.0
; SEQ. ID NO. 2 PatentIn Ver. 2.0
; LENGTH: 483
; TYPE: PRF
; ORGANISM: Muba sp.
; US-08-833-553-2

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Query Match	38.8%;	Score 965.5;	DB 3;	Length 483;
Best Local Similarity	42.9%;	Pred. No. 6.1e-90;		
Matches 204;	Conservative	75;	Mismatches 180;	Indels 17;
				Gaps 10;

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QY      11 LPLKIPGGGIFPPGLGRIKORIDIFFQRDERFKNTIKNSIVFKNMPEQ-F-EISS 68
        |::|||::|||::|||::|||::|||::|||::|||::|||::|||:
Db      16 LPTRIPGGYGPLVGPLKDRLDYFMFGQEPETFRSMATKSTVFRTMPPTPFPGGV 75
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Db 76 DPRVATLDSLSALFDLEVEYKRNILIGDIMPISLFTGSDTRVVVYLDPSPEDHAAVKS 135

[illegible]

Db 195 DPSVSPDVGENGFVMDKMLALQLLPYKVG--AIPOPLEELHLSPLPFLVSRDVRK 252
 QY 247 IYEAFSSSSGSFLPDEAF-KQGI DREKACRNLPVLGNAGVAGMKVLPFLTKVGVGTACED 305

Db 253 LYEPEVKQCGEVRRARTEHGLSKDAINNIILFVLGNAGFGSPVFPPTLLTTIGRDTG 312

Cy 306 LHRRLAEVHTTYKEGCL--TFSLERKSLKSVVEALRIEPPVPFGAKEDIIVQ 363

Db 313 LREKLKDEYVRVWKKSRDEHKRPSFETVREKELVSTVYEVLNLNPPVLQYGRARTDTPLN 372

Qy 364 SHDSFPIKKGGTITFGYQPFATKDKPIFKDSEKFGDGRFVGBEGEKLKYVWVSNERTV 423

DB 373 SHDAAFKVEGSEBLLCGYPLWKRDPVAFDDPELFAFERFMG-SUKELLKLVFMSNPEIC 433

QY 424 EPTAENKOCGRKULVVLIGRIWVEFFLLSYDTFTVEADLPLGPAVKESLRTATD 479

RESULT 4

; Sequence 2, Application US/09418222
 ; Patent No. 623898
 ; GENERAL INFORMATION:
 ; ADDICANT, Hansler Alay

```

; APPLICANT: Lersch, Konrad
;
; APPLICANT: Muehl, Andreas
;
; APPLICANT: Silke, Natasha
;
; TITLE OF INVENTION: HYDROPEROXIDE LYASES

```

FILE REFERENCE: Hydroperoxide, lysase
 CURRENT APPLICATION NUMBER: US/09/418,222
 CURRENT FILING DATE: 1999-10-14
 EARLIER APPLICATION NUMBER: US/09/833,553
 EARLIER FILING DATE: 1997-04-07
 NUMBER OF SEQ ID NOS: 11
 SOFTWARE: Patent Ver. 2.0
 SEQ ID NO 2
 LENGTH: 483
 TYPE: PR
 ORGANISM: Musa sp.
 US-09-418-222-2

Query Match 38.8%; Score 965.5; DB 3; Length 483;
 Best Local Similarity 42.9%; Pred. No. 6; Le 90;
 Matches 204; Conservative 75; Mismatches 180; Indels 17; Gaps 10;

11 LPLKPIGCGYPPFLGPIKDRDYFYFGQDEFFRSRTYKNSIVFRANMPG-P-FISS 68
 16 LPTKPIGCGYPPFLGPIKDRDYFYFGQDEFFRSRTYKNSIVFRANMPG-P-FISS 75
 69 DSRVVLADLSPFLPPTAKYKRNILDTWPSLSFTGNTKCAVLDSETHSVLR 128
 76 DSRVVLADLSPFLPPTAKYKRNILDTWPSLSFTGNTKCAVLDSETHSVLR 135
 129 LPLASRHRDRIPLFRSSLSSEFVKLEEDLSSEKKIADPNISIDSMSPDYVRL--S 186
 136 FCLBLRGAKTWSSPLSLDVMALTIQGIAGOSAGLFGPQKCI-FAFLCKSIIGA 194
 187 DCPDPSKLAAGCPQFMDLTVFQALPLASIGLPIFSVPEDEVHTIPLPFPVKSQYRK 246
 195 DSVSPDVGNGFVMDLQMLALQILPTVKG--ALPQLEHLLHSFPLPPLVSRDYRK 252
 247 LYEAFSSSGSFIDEAE-KQIDREKACHNVLGAFNAYGGMVLEPTLLKAVGTAGSD 305
 253 LYEAFSSSGSFIDEAE-KQIDREKACHNVLGAFNAYGGMVLEPTLLKAVGTAGSD 312
 306 LHRKLAERTTYVEEGSL--TFSALEKNSILKSVTEALRIEPPVPOYGAKEDEVIO 363
 313 LHRKLAERTTYVEEGSL--TFSALEKNSILKSVTEALRIEPPVPOYGAKEDEVIO 372
 364 SHDSSEKIKKGGFTIPGQOPATKDKIPFDSSEKFGVGRVGESEKLLKYVNSNERETV 423
 373 SHDAAFVKKGGELCGQPLVMBDPAVDFDPTFAPRFPWG-SGKELLKYVNSNGPETG 431
 424 EPTANMOCPCGNLVVLIIGRIWVFFLYVTFVAVDLPLGPAVFKSLTRATD 479
 432 TPTANMOCADYVETACILMAEIFYRYDEV--CAD---DAISVTKLDRAE 481

RESULT 5
 US-09-537-357-33
 Sequence 33; Application US/09537357
 Sequence 33; Application US/09537357
 GENERAL INFORMATION:
 APPLICANT: Alan Braash
 APPLICANT: Nathalie Tijet
 TITLE OF INVENTION: MUSKRELON (CUCUMIS MELO) HYDROPEROXIDE
 FILE REFERENCE: 06027.0002
 CURRENT APPLICATION NUMBER: US/09/537,357
 CURRENT FILING DATE: 2000-03-29
 NUMBER OF SEQ ID NOS: 56
 SOFTWARE: Patented for Windows Version 4.0
 SEQ ID NO 38
 LENGTH: 483
 TYPE: PR
 ORGANISM: Banana
 US-09-537-357-33

Query Match 38.8%; Score 965.5; DB 3; Length 483;
 Best Local Similarity 42.9%; Pred. No. 6; Le 90;
 Matches 204; Conservative 75; Mismatches 180; Indels 17; Gaps 10;

11 LPLKPIGCGYPPFLGPIKDRDYFYFGQDEFFRSRTYKNSIVFRANMPG-P-FISS 68
 16 LPTKPIGCGYPPFLGPIKDRDYFYFGQDEFFRSRTYKNSIVFRANMPG-P-FISS 75
 69 DSRVVLADLSPFLPPTAKYKRNILDTWPSLSFTGNTKCAVLDSETHSVLR 128
 76 DSRVVLADLSPFLPPTAKYKRNILDTWPSLSFTGNTKCAVLDSETHSVLR 135
 129 LPLASRHRDRIPLFRSSLSSEFVKLEEDLSSEKKIADPNISIDSMSPDYVRL--S 186
 136 FCLBLRGAKTWSSPLSLDVMALTIQGIAGOSAGLFGPQKCI-FAFLCKSIIGA 194
 187 DCPDPSKLAAGCPQFMDLTVFQALPLASIGLPIFSVPEDEVHTIPLPFPVKSQYRK 246
 195 DSVSPDVGNGFVMDLQMLALQILPTVKG--ALPQLEHLLHSFPLPPLVSRDYRK 252
 247 LYEAFSSSGSFIDEAE-KQIDREKACHNVLGAFNAYGGMVLEPTLLKAVGTAGSD 305
 253 LYEAFSSSGSFIDEAE-KQIDREKACHNVLGAFNAYGGMVLEPTLLKAVGTAGSD 312
 306 LHRKLAERTTYVEEGSL--TFSALEKNSILKSVTEALRIEPPVPOYGAKEDEVIO 363
 313 LHRKLAERTTYVEEGSL--TFSALEKNSILKSVTEALRIEPPVPOYGAKEDEVIO 372
 364 SHDSSEKIKKGGFTIPGQOPATKDKIPFDSSEKFGVGRVGESEKLLKYVNSNERETV 423
 373 SHDAAFVKKGGELCGQPLVMBDPAVDFDPTFAPRFPWG-SGKELLKYVNSNGPETG 431
 424 EPTANMOCPCGNLVVLIIGRIWVFFLYVTFVAVDLPLGPAVFKSLTRATD 479
 432 TPTANMOCADYVETACILMAEIFYRYDEV--CAD---DAISVTKLDRAE 481

RESULT 6
 US-09-078-173A-26
 Sequence 26; Application US/09078173A
 Sequence 26; Application US/09078173A
 GENERAL INFORMATION:
 APPLICANT: Alan Braash
 APPLICANT: Ian M. Whitehead
 APPLICANT: Alan Siuarengo
 APPLICANT: Alan Siuarengo
 APPLICANT: Alan Siuarengo
 APPLICANT: Alan Siuarengo
 TITLE OF INVENTION: GUAVA (PSIDIUM GUJAVANA) 11-HYDROPEROXIDE
 FILE REFERENCE: 06027.0001
 CURRENT APPLICATION NUMBER: US/09/078,173A
 CURRENT FILING DATE: 1998-05-13
 NUMBER OF SEQ ID NOS: 127
 SOFTWARE: Patented for Windows Version 3.0
 SEQ ID NO 26
 LENGTH: 483
 TYPE: PR
 ORGANISM: Musa sp. (banana)
 US-09-078-173A-26

Query Match 38.8%; Score 961.5; DB 3; Length 483;
 Best Local Similarity 42.9%; Pred. No. 1.6e-89;
 Matches 204; Conservative 74; Mismatches 181; Indels 17; Gaps 10;

11 LPLKPIGCGYPPFLGPIKDRDYFYFGQDEFFRSRTYKNSIVFRANMPG-P-FISS 68
 16 LPTKPIGCGYPPFLGPIKDRDYFYFGQDEFFRSRTYKNSIVFRANMPG-P-FISS 75
 69 DSRVVLADLSPFLPPTAKYKRNILDTWPSLSFTGNTKCAVLDSETHSVLR 128
 76 DSRVVLADLSPFLPPTAKYKRNILDTWPSLSFTGNTKCAVLDSETHSVLR 135
 129 LPLASRHRDRIPLFRSSLSSEFVKLEEDLSSEKKIADPNISIDSMSPDYVRL--S 186
 136 FCLBLRGAKTWSSPLSLDVMALTIQGIAGOSAGLFGPQKCI-FAFLCKSIIGA 194
 187 DCPDPSKLAAGCPQFMDLTVFQALPLASIGLPIFSVPEDEVHTIPLPFPVKSQYRK 246
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 364 SHDSSEKIKKGGFTIPGQOPATKDKIPFDSSEKFGVGRVGESEKLLKYVNSNERETV 423
 373 SHDAAFVKKGGELCGQPLVMBDPAVDFDPTFAPRFPWG-SGKELLKYVNSNGPETG 431
 424 EPTANMOCPCGNLVVLIIGRIWVFFLYVTFVAVDLPLGPAVFKSLTRATD 479
 432 TPTANMOCADYVETACILMAEIFYRYDEV--CAD---DAISVTKLDRAE 481

187 DGTDSKLAABGQFDMVLVFOALPLASIGLPIKISVEEDLVHTITPLPPVSGYRK 246
195 DESVSPDVGNGVMDKMLALQULPTVKVG--AIPQLESIILHSPPLPPLVSDNYRK 252
247 LYRPAFYSSSSFLDEAB--KQIDBERACAHVPLAGRNKQCKVPTLLKMTGKASD 305
253 LYEPVKKQGOVVRABSTHSLSDHAINIILPVAGRNKQSGSVPTLLTIGDCKTG 312
306 LHRKLAERVITVYKEOGL--TPSALBEMSLKSVYVLEALRIBPPVPOYGAKEDIYIO 363
313 LREKLMDEVRRVWKSREGRSPFVTRMELVRSVYVLEALRIBPPVPOYGAKEDPTLN 372
364 SHDSFKIKKGETTIGYOPATNDPKIKQSEKRVGDRFVGEGBEKLKYYMSNERETV 423
373 SHDAAKVGKGBLLCGYQPLVNRDPAVDDPETPAPRPMG--SKCKLAKYVMSNGSTG 431
424 BPTSNKQCGKMYLIGRIMVVEPFLAKYPTVTEALDPAVKNKSLTRATD 479
432 TPTANKQCAADYVETACLMALIFRIDEV--CND---DALSVTLDBABE 481

RESULT 7
US-10-042-991-26
Sequence 26, Application US/10042991
Patent No. 6780621
GENERAL INFORMATION:
APPLICANT: Ian W. Whitehead
INVENTOR: Alan Stuartenko
APPLICANT: Duncan Gaekin
APPLICANT: Alan Stuartenko
APPLICANT: Nathalia Tillet
TITLE OF INVENTION: GUAVA (PSIDIUM GUJAVARA) 13-HYDROPEROXIDE
FILE REFERENCE: 06027.0001UD
CURRENT APPLICATION NUMBER: US/10/042, 991
CURRENT FILING DATE: 2002-01-09
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 26
LENGTH: 483
TYPE: EXT
ORGANISM: Guava sp. (banana)
US-10-042-991-26

Query Match 38.64; Score 961.5; DB 4; Length 483;
Best Local Similarity 42.94; Pred. No. 1.6e-89;
Matches 204; Conservative 74; Mismatches 181; Indels 17; Gaps 10;

11 LPLKPIPGYGPPLGPIKDXRDVFFQSDHFPFASRIKTNSTVFRANMPG--P-FISS 68
16 LPTPIPGSYGPPVLGDLRIDYTFQGPETFPFASMAHKSIVKTNAPVPPPPGV 75
69 DSNVYVLDLALSPILITPAKTKENIILDTGTPSLSFTGNIRTCAYLIDSTETSVYLR 128
76 DRYVYVLDCTSPSALDLEVERKQILIDMNSLSFTGDIKVVYVLDSPBDKAVKS 135
129 LFLSPLASRRDRITLFRSSLSBMYKXEDKCSBKXKIADNSISDSMDVYRLL--S 186
136 FCLLELRGAKVWSSFLSNLDWALATIEOGLAKQSGAGLFGPLQKCI--PAFLCKSTIGA 194
187 DGTDSKLAABGQFDMVLVFOALPLASIGLPIKISVEEDLVHTITPLPPVSGYRK 246
195 DESVSPDVGNGVMDKMLALQULPTVKVG--AIPQLESIILHSPPLPPLVSDNYRK 252
247 LYRPAFYSSSSFLDEAB--KQIDBERACAHVPLAGRNKQCKVPTLLKMTGKASD 305
253 LYEPVKKQGOVVRABSTHSLSDHAINIILPVAGRNKQSGSVPTLLTIGDCKTG 312
306 LHRKLAERVITVYKEOGL--TPSALBEMSLKSVYVLEALRIBPPVPOYGAKEDIYIO 363
313 LREKLMDEVRRVWKSREGRSPFVTRMELVRSVYVLEALRIBPPVPOYGAKEDPTLN 372
364 SHDSFKIKKGETTIGYOPATNDPKIKQSEKRVGDRFVGEGBEKLKYYMSNERETV 423

373 SHDAAKVGKGBLLCGYQPLVNRDPAVDDPETPAPRPMG--SKCKLAKYVMSNGSTG 431
424 BPTSNKQCGKMYLIGRIMVVEPFLAKYPTVTEALDPAVKNKSLTRATD 479
432 TPTANKQCAADYVETACLMALIFRIDEV--CND---DALSVTLDBABE 481

RESULT 8
US-09-078-173A-25
Sequence 25, Application US/09078173A
Patent No. 6200794
GENERAL INFORMATION:
APPLICANT: Ian W. Whitehead
INVENTOR: Alan Stuartenko
APPLICANT: Duncan Gaekin
APPLICANT: Alan Stuartenko
APPLICANT: Nathalia Tillet
TITLE OF INVENTION: GUAVA (PSIDIUM GUJAVARA) 13-HYDROPEROXIDE
FILE REFERENCE: 06027.0001
CURRENT APPLICATION NUMBER: US/09/078, 173A
CURRENT FILING DATE: 1998-05-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 25
LENGTH: 480
TYPE: EXT
ORGANISM: Capsicum annuum (green pepper)
US-09-078-173A-25

Query Match 37.14; Score 922.5; DB 3; Length 480;
Best Local Similarity 40.14; Pred. No. 1.5e-85;
Matches 189; Conservative 92; Mismatches 171; Indels 19; Gaps 7;

2 ATFFSSSPPLPLKPIPGYGPPLGPIKDXRDVFFQSDHFPFASRIKTNSTVFRANMPG 61
13 ATPLS---LPTVKIPGSYGPPLGDLRIDYTFQGPETFPFASMAHKSIVKTNAPVPPPPGV 68
62 P--GPISSDSRVYVLDLALSPILITPAKTKENIILDTGTPSLSFTGNIRTCAYLIDPS 119
69 PCPPFLGVRNVAVLDVKSALHDMETVFNKAVNVLDGPPSVYVGDGRVCAVYDPS 128
120 ETEHSVYKELFLSPLASRRDRITLFRSSLSBMYKXEDKCSBKXKIADNSISDSM--SF 176
129 EKHQIQKMFSLDIKSSKTVVTLWEIDTLFGTRESPLSKSKASLPLALQKFLNFR 188
119 DYVRLISDGTPODSKLAABGQFDMVLVFOALPLASIGLPIKISVEEDLVHTITPLPP 238
189 FSLTFLGADPPSAPSPISNGSFAYLDMLALQULPTVKSIG--VLPQLESIIVHVSFSYPR 245
239 PVKSGTRKIVBARYSSGSLDEABEQ--GIDREKACAHVPLAGRNKQCKVPTLLK 297
246 LVRGAGBGLAHKLAERVITVYKEG---LTPSALBEMSLKSVYVLEALRIBPPVPO 352
298 LVRGAGBGLAHKLAERVITVYKEG---LTPSALBEMSLKSVYVLEALRIBPPVPO 352
305 --GIDREKACAHVPLAGRNKQCKVPTLLKMTGKASD 363
353 YKAKAKEDVIOSSHDSFKIKKGETTIGYOPATNDPKIKQSEKRVGDRFVGEGBEKLK 412
363 YKAKAKEDVIOSSHDSFKIKKGETTIGYOPATNDPKIKQSEKRVGDRFVGEGBEKLK 422
413 YVMSNERETVPTANKQCAADYVETACLMALIFRIDEV--CND---DALSVTLDBABE 463
423 YVMSNERETVPTANKQCAADYVETACLMALIFRIDEV--CND---DALSVTLDBABE 473

RESULT 9
US-09-537-357-32
Sequence 32, Application US/09537357
Patent No. 6271016
GENERAL INFORMATION:

```

1  APPLICANT: Alan Brash
2  APPLICANT: Nathalie Tjelt
3  TITLE OF INVENTION: MUSHMELON (CUCUMIS MELO) HYDROPEROXIDE
4  TITLE OF INVENTION: LYASE AND USSS THERAP
5  FILE REFERENCE: 06027.0002
6  CURRENT APPLICATION NUMBER: US/09/537.357
7  CURRENT FILING DATE: 2000-03-25
8  NUMBER OF SEQ ID NOS: 56
9  SOFTWARE: FastSeq for Windows Version 4.0
10 SEQ ID NO 32
11 LENGTH: 480
12
13 TYPE: PRT
14 ORGANISM: Capsicum annuum (green pepper)
15 US-09-537-357-32

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Query Match	37.1%	Score 922.5;	DB 3;	Length 480;
Best Local Similarity	40.1%	Pred. No. 1.5e-85;		
Matches 189; Conservative	92;	Mismatches 171;		
		Indels 19;	Gaps 7	

Qy 2 ATSSSSSHFLPLAKPIPGYGFPELAPIKDRIVYPIRQGNDEFFRSRLTKTKNTVTPAANP 61
13 ATATIS-----IPKRIKPGSGPLPGLGMDLNTMNVQGLPFGSGVKNSTSTYKRNK 68
Qy 62 P--GPIISNSNVTVLDAISFLPIEDMANIKRNLIDQITWYSLSTKTKITCAVDIS 119

DB 69 PCPPFFLGVPNNVAVLDVKSFAHLFDMETKANKLVGDPMPSVYTTGDMRCVCAVLDTS 128

OV 120 ETEHSVLRKFLPLASRHHDRPIPLERSLSLMPVXLJEDKLSKKKIIADFNISDSM-6F 178

Db 129 EPKHQIQKNSPLDILKRSSKTVPTLVKEIDPLFGFESDLSKSKASILPALKQFLNF 188

Db 189 FSLTFLGADPASPRIANGFAYLDALAIQAPFVSIG---VLQPLEIIFVHSFSVYP 245

Dh
Qy

239 PVKSGRKLTAEAFYSSSGSLDDEAKG-CIDREKACGNLVPLAGFNAYGGMKVLPFLTK 297

246 LVRGVEKLIRKVKSEAKKEVITRAQDFPOLTEOEAIHNLPLFGFNAFGPTILPFLPI- 304

298 WGTAGEDHHRKLAEBVRYTYKEGG-----LTFSALBKMSLLKSVYBALRIBPPVPFO 352

DB 305 - - GNLCJDBNANMFBKLRABVBAVG IINGENUSF BSVYBNBIAU QGZ * L DBNACDZ E Z * V QZ

OY 353 YOKAKEDIVIOSHDSFKIKKGETTIFGYOPATDKPKI PKXSEKPVGD RFVEEGBKLL 412

Db 363 YARAKDPMLSHSDVYEIKKGEILCGYQPLVMKDPKYFDEPEKFMLEPFTKEKGKELLN 422

Db 423 YLFMSNGPOTGSPTESNKQCAAKDAVLTASLIVAYTEQKYDSVSFSGSL 473

RESULT 10
US-10-042-991-25
10/1/2004 10:00:00

Patent No. 6780621
; GENERAL INFORMATION:
; APPLICANT: Ian M. Whitehead

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: APPLICANT: Alan Sissarenko
: APPLICANT: Duncan Gaskin
: APPLICANT: Alan Brash

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: CURRENT APPLICATION NUMBER: US/10/042,991
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: CURRENT PILING DATE: 2002-01-09
:
: NUMBER OF SEQ ID NOS: 27
:

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: SOFTWARE:FastSeq for Windows Version 3.0
: SEQ ID NO 25
: LENGTH: 480

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ORGANISM: Capsicum annuum (green pepper)

US-10-042-991-25

Query Match	37.1%	Score 922.5;	DB 4;	Length 480;
Best Local Similarity	40.1%;	Pred. No. 1.5e-85;		
Matches 189;	Conservative 92;	Mismatches 171;	Indels 19;	Gaps 7;

2 ATPSSSSDELPLKPIPGCGPFLPGIKORYDYFEGQRDEFFRSRTTKYNSVTFRANP 61
13 ATPIS-----LPVRIKPGSYGFPLLGCLMELDYNNFQKLDPFSKREKYNSTVFRINP 68

62 P--GPTISDSRVVLLDALSPILLEDTAKVEKRNILDTYMPSPSTGNIRTCAYLDPS 119
| : : | | | | | : | : : : : : : : | | | |
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120 ETEHSVLRKRLPTSLASRHRDPIPLFRSSLSSEMFVTEDEKLSSEKKKLDNFNSISDSM-SF 178

129 EPKHTQIQNPSLIDILKRSSKTWPVLVKELDTLFGTPESSDLSKSKSALLPALQKFLPNF 188

179 DYVFRLLSDGTPDSKTLAEGPGMFEDLWLVQDLAPLASIGLPKIRSVPEEDLVIHITPLPFF 238

189 PSILTFGLGADPSASPEIANGSFAYLDWLAIQLAPTSIG--VLQPLEEIFVHSFSYPYF 245

246 LVRGGYEKLKPVTSSEAKSVLTRAQDFQLTEQAHHLLPILGPNACGGFTILEPTLL- 304

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298 WGTAGEDLHKLAEEVRTTVKEEG-----LTPSALSKMLSLKSVYEAALRIEPPVPFQ 352
      |:::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|
305 --GNLGDENAEOMEKLRKEVREKVGVTNOENLSFESYKEMELVQSFYVYESTRLSPVPSQ 362

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353 YGAKAKEDIVIQSHDSFKIKGGTTCYOPATKDPKIFKDSKPFVDRFVGECEKLLK 412

413 YYYWSNERETVEPTAENKQCPGNLVYLIGRIWVVEFFLYIDPTFVAVDL 463

423 YLIPNSGPGQSPTESNKQCAKDAVTLTASLIIVAIYIFQKIDSVSFSSGS 473

RESULT 11
US-09-078-173A-2
Sequence 2, Application US/09078173A

Patent No. 6200794
GENERAL INFORMATION:
APPLICANT: Ian M. Whitehead

APPLICANT: Duncan Gaskins
APPLICANT: Alan Brash
APPLICANT: Nathalie Tillet

TITLE OF INVENTION: GUAVA (PSIDIUM GUAJAVA) 13-HYDROPEROXIDE
TITLE OF INVENTION: LYASE AND USES THEREOF
FILE REFERENCE: 06027.0001

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SEQ ID NO 2
LENGTH: 476
TYPR. PPT

ORGANISM: *Psidium Guajava* (guava)
US-09-078-173A-2

Query Match	56.43	Score 504	26.5	Length 170
Best Local Similarity	41.94	Pred. No. 2.4e-83		
Matches 198; Conservative	79;	Mismatches 163;	Indels 32;	Gaps 12

QY

1 MATPSSSSP-ELPLKPIPGGYGPPGLGIKDRIYIFYFCORDEFFPSRITKTNSTVFRAN 59
::|||::|||::|||::|||::|||::|||::|||::|||::|||::|||::|||::|||::|||
9 LSPSSDRPTTLPVRTIPGSYGMPILGPISDRLDYFWFOCPETPFPRKRIEKKSTVFRAN 68

60. MPP-GPEISS-DSRVVLLDALSPILPPTAKVEKNIIDGTYMPSISFTGNIRTCAYLD 117

09 VERCFETSNVENVVYUCCUGI NUBH DNUA *BMON*ZD OCT 68

Query Match	36.2%	Score 901,	Id 4,	Length 476,
Best Local Similarity	41.3%	Pred. No. 2,46-33,		
Matches	198	Conservative	79,	Mismatches 163, Indels 33, Gaps 152
Qy	1	MATRRSSP-RLEPKIQGQYCPKPLKIDYKYYGKGGPFRITNGTNGRYAN		
Dy	1	LSPSSRPRLPLPRITIDYSGWPLPISDLRLDYMGGPFRIRKIKRYKATYFRAN		
Qy	60	MP-QPISG-DSRVVVLIDLSPFLPTPAVAKKNILIDTWSLSPFGNITCAIYD		
Dy	60	VPPCPFFSNVAPVAVVVLIDCSFPAHDPMEIKRSNVAVDPMSSYKRYGNIHCAYLD		
Qy	118	PSTHTYVAKRLPLASTLSRHRDPLPFRSLISMYPLKEDKLSKKGLADNDSIGMS		
Dy	123	TSEPHQAVNADNKLAKSSKRWSEVSNLDMDVDEESLAK---DQNN---SVI		
Qy	178	P--DYVFRLLS---DGTDFSKLAAGCGQMTLMLPPLAPLASIGLPIRVSFEEL		
Dy	181	PLPLQGLFRLSKSITIGDPLAPLQVMAKSSMLMDKRLDMLPLTNG--VLQDPVEI		
Qy	223	VITHTLPAPFPYSGRGLKLYAVSAGSFLEBAG--GIDRKKCNVLPFLAPNAYG		
Dy	238	FLISNAVAPVAVDSDNKLVOFTEKGGSEKVRKKAPFLHQBHINALLPLPAPNACG		

Query Match	36.2%	Score 901;	DB 3;	Length 480;
Best Local Similarity	41.9%	Pred. No. 2; 46-83;		
Matches	198;	Conservative	79;	Mismatches 169; Indels 32; Gaps 12;
Qy	1	MATPSSSP-ELPKIRIGGYGPPLPIKORVDYFFQSGDEPSRRTTKTNSVTPKAN	59	
Dh	13	LSBSSSRRLPTKPKVILKPSGHWLPSLSDLVYVLTQGTFFPKTKERKNSVTPKAN	72	
Oy	60	MPP-GRPIS-DSNWWLTLASPIPLPTKAKKKNLITQGTFFNSLSTNGITATVLD	117	
Dh	73	VPCPFPFSSNPVVVVVWLDCSPFHLFEMELYEKSNVLDGPFMSVKTNGIRIVATLD	132	
Oy	118	PSTSTHSEKRLPLSLASLRHRBPLPLFSSLSSEKVLDEKLSKCKKLAAPNISDSMS	177	
Dh	133	TSPHQAQVKNNAADLLKSSKSWSESVISNLDMWDTIESLAL-DBAN--SVI	184	
Oy	178	F---DYRSLA-----DTPSLKLAGGPGPLPVYVQLALGLPKRIFSVEDL	228	
Dh	185	PLQGLPRLPMSLISITGADPRASPVAKSGVAMLDMAQLPLTINIG--VGLQVRI	241	
Oy	229	VHITPLPFPFVPSGRVLYEATYSSSGSLPDEAKQ-GIDREKCKNVPLADPAAVQ	287	
Dh	242	PLMSAAVPAVLASVDNGLVQPLFEKERGAVERAKKAPGLTIOBZHLINLLPLGPAAGG	301	
Oy	288	MKYLEPTLLKAVGTAGIEDLRKLAEEVRTVYBERG-LTFSNLEKNSLLSVYEAIRI	345	
Dh	302	FSTPLPFLSLNISLSDTQGRRLKREYR-----KGPALASVAMENELASVSTLRL	357	
Oy	346	BPPVPGQVQVLTQSDHSPKPKIKGTIGTQVPPAPAKPKIPROSEKAPPPGPR	405	
Dh	358	NPPVFPVPAARNDPOLASHDHSVDFVKKKELLGQVXWVTHHPLKVDDEPSNSDRV-	415	
Oy	406	EGRKLLKCYVMSNREYVETPLAMNQPGKQILVVLGIVWVEPLLYVTPL	457	
Dh	416	QNSLLDLYVMSNDPTGTTPENNQQAKQYVTLTALCLFAVAFRRYSVY	467	

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RESULT 14
US-10-042-991-3
Sequence 3, Application US/10042991
Patent No. 6786621
GENERAL INFORMATION:
APPLICANT: Ian M. Whitehead
APPLICANT: Alan Sinaerenko
APPLICANT: Duncan Gaekin
APPLICANT: Alan Braeh
APPLICANT: Nathalie Tije
TITLE OF INVENTION: GUAVA (PSIDIUM GUAJABA) 13-HYDROPEROXIDE
TITLE OF INVENTION: LYSINE AND USES THEREOF
CURRENT FILING DATE: 2002-01-09
CURRENT PILING DATE: 2002-01-09
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 3
LENGTH: 480
TYPE: PR
ORGANISM: Psidium Guajava (guava)
US-10-042-991-3

Query Match 36.2% Score 901; DB 4; Length 480;
Best Local Similarity 41.9%; Pred. No. 2.5e-83;
Matches 198; Conservative 79; Mismatches 163; Indels 32; Gaps 12;

QY 1 MATPSSASP-ELPLKPIGCGYGPFLGPIKORVDYFYQGRDEPFRSRTTKNSTVFRAN 59
DB 13 LSPSSPPTTLTPVPTIGSYGMPLGPISDRLDYFMFGPPTFRKRIKRYSTVFRAN 72
QY 60 MPP-GPFISS-DSRVVLLDALSPFLPFTAKYERKNIIDGTWPSLSFTGNIRTCAYLD 117
DB 73 VPCPPEPSSNNVNNVYVLDCEPFAHLPMEIYKSNVAVGDPMPBVKIKGNIRTCAYLD 132
QY 118 PSTERSYKXLEPLSLASHDRFIPLRSSLSSEMFVYKLEKLSKKKIDAPNSISDSNS 177
DB 133 TSPQHAQVKNPMDILKSSSVESVYISLDTWMTISSLAK-----DGNH---SVI 187
QY 178 F---DYVPLLS-----DQTPDSKLAEGPGFDMVLPOLAPLASIGLPKISVFEOL 228
DB 185 PLOKPLFPLSKSIIGADPAASQVAKSGYMLDRMLALQLPTINIG--VLOPLVEI 241
QY 229 VHTIPLPEFPVNSGYRKYEAFFSSSSGFIDBAEKQ-GIDREKACHNVLPLAGFNAYG 287
DB 242 FLHSMVAFPALVSGDYNKYOTIEKGRVAYEAKQERGTQKALHMLPILIGFNAPG 301
QY 288 MKVLPTLLKAVGTAGDHLRKLAEVPTVKEEG--LTPSALERKSLKSYVEALRI 345
DB 302 PSILPLTLLSNILSDTGTGLDRLKEVRA---KGGPALSASVYKEMELVSVYETLRL 357
QY 346 EPPVPOYKAKEDIVTOSHDSFPKIKGTITIGYQPPATKDPKIPKDESEKVPVGE 405
DB 358 NPVPFOYARAKDPOLKSHDSVPVYKGBELGQYQKVMWTDPKVPDESPNSDRFY-- 415
QY 406 EGEKLLKCYVNSNERETVEPTAKNCCPGKRLVYLIGIMVVEPFLAYDTFT 457
DB 416 QNSLDDVLYMSNGPQTGTFTSSNCKAANDYVTLACLFLVAVMFRKINSVT 467

RESULT 15
US-09-078-173A-4
Sequence 4, Application US/09078173A
Patent No. 6280734
GENERAL INFORMATION:
APPLICANT: Ian M. Whitehead
APPLICANT: Alan Sinaerenko
APPLICANT: Duncan Gaekin
APPLICANT: Alan Braeh
APPLICANT: Nathalie Tije
TITLE OF INVENTION: GUAVA (PSIDIUM GUAJABA) 13-HYDROPEROXIDE
FILE REFERENCE: 06027.000
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CURRENT APPLICATION NUMBER: US/09/078,173A
CURRENT FILING DATE: 1998-05-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 483
TYPE: PR
ORGANISM: Psidium Guajava (guava)
US-09-078-173A-4

Query Match 36.2% Score 901; DB 3; Length 483;
Best Local Similarity 41.9%; Pred. No. 2.5e-83;
Matches 198; Conservative 79; Mismatches 163; Indels 32; Gaps 12;

QY 1 MATPSSASP-ELPLKPIGCGYGPFLGPIKORVDYFYQGRDEPFRSRTTKNSTVFRAN 59
DB 16 LSPSSPPTTLTPVPTIGSYGMPLGPISDRLDYFMFGPPTFRKRIKRYSTVFRAN 75
QY 60 MPP-GPFISS-DSRVVLLDALSPFLPFTAKYERKNIIDGTWPSLSFTGNIRTCAYLD 117
DB 76 VPCPPEPSSNNVNNVYVLDCEPFAHLPMEIYKSNVAVGDPMPBVKIKGNIRTCAYLD 135
QY 118 PSTERSYKXLEPLSLASHDRFIPLRSSLSSEMFVYKLEKLSKKKIDAPNSISDSNS 177
DB 136 TSPQHAQVKNPMDILKSSSVESVYISLDTWMTISSLAK-----DGNH---SVI 187
QY 178 F---DYVPLLS-----DQTPDSKLAEGPGFDMVLPOLAPLASIGLPKISVFEOL 228
DB 188 PLOKPLFPLSKSIIGADPAASQVAKSGYMLDRMLALQLPTINIG--VLOPLVEI 244
QY 229 VHTIPLPEFPVNSGYRKYEAFFSSSSGFIDBAEKQ-GIDREKACHNVLPLAGFNAYG 287
DB 245 FLHSMVAFPALVSGDYNKYOTIEKGRVAYEAKQERGTQKALHMLPILIGFNAPG 304
QY 288 MKVLPTLLKAVGTAGDHLRKLAEVPTVKEEG--LTPSALERKSLKSYVEALRI 345
DB 305 PSILPLTLLSNILSDTGTGLDRLKEVRA---KGGPALSASVYKEMELVSVYETLRL 360
QY 346 EPPVPOYKAKEDIVTOSHDSFPKIKGTITIGYQPPATKDPKIPKDESEKVPVGE 405
DB 361 NPVPFOYARAKDPOLKSHDSVPVYKGBELGQYQKVMWTDPKVPDESPNSDRFY-- 418
QY 406 EGEKLLKCYVNSNERETVEPTAKNCCPGKRLVYLIGIMVVEPFLAYDTFT 457
DB 419 QNSLDDVLYMSNGPQTGTFTSSNCKAANDYVTLACLFLVAVMFRKINSVT 470

Search completed: October 5, 2005, 07:00:26
Job time : 46 secs
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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: October 5, 2005, 07:00:33 ; Search time 168 Seconds
(without alignments)
1188,517 Million cell updates/sec

Title: US-09-884-260A-7
Perfect score: 2488SSSPPLPLKIPGCT.....DPLDAPNVEKSLTRATNY 481
Sequence: 1 MATPSSSSPPLPLKIPGCT.....DPLDAPNVEKSLTRATNY 481

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1846076 seqs, 41511600 residues

Total number of hits satisfying chosen parameters: 1846076

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 01
Maximum Match 1004
Selecting first 45 summaries

Database: Published Applications AA:*

- 1: /cgn2_6/prodate/1/pubpaa/US07_PUBCOMB.pep.*
- 2: /cgn2_6/prodate/1/pubpaa/US07_PUBCOMB.pep.*
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- 5: /cgn2_6/prodate/1/pubpaa/US07_PUBCOMB.pep.*
- 6: /cgn2_6/prodate/1/pubpaa/US07_PUBCOMB.pep.*
- 7: /cgn2_6/prodate/1/pubpaa/US08_PUBCOMB.pep.*
- 8: /cgn2_6/prodate/1/pubpaa/US08_PUBCOMB.pep.*
- 9: /cgn2_6/prodate/1/pubpaa/US09_PUBCOMB.pep.*
- 10: /cgn2_6/prodate/1/pubpaa/US09_PUBCOMB.pep.*
- 11: /cgn2_6/prodate/1/pubpaa/US09_PUBCOMB.pep.*
- 12: /cgn2_6/prodate/1/pubpaa/US09_PUBCOMB.pep.*
- 13: /cgn2_6/prodate/1/pubpaa/US10_PUBCOMB.pep.*
- 14: /cgn2_6/prodate/1/pubpaa/US10_PUBCOMB.pep.*
- 15: /cgn2_6/prodate/1/pubpaa/US10_PUBCOMB.pep.*
- 16: /cgn2_6/prodate/1/pubpaa/US10_PUBCOMB.pep.*
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- 22: /cgn2_6/prodate/1/pubpaa/US10_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2488	100.0	481	US-09-884-260A-7	Sequence 7, Appl1
2	2488	99.8	481	US-10-434-991-9	Sequence 9, Appl1
3	2483	99.0	487	US-09-884-260A-15	Sequence 15, Appl1
4	2339	94.0	483	US-10-434-991-10	Sequence 10, Appl1
5	1613.5	64.9	483	US-10-686-947-156	Sequence 156, Appl1
6	1560	62.7	487	US-10-434-991-6	Sequence 6, Appl1
7	1560	62.7	522	US-10-424-599-203458	Sequence 203458, Appl1
8	1546.5	62.2	492	US-10-425-114-40766	Sequence 40766, A
9	1546.5	62.2	492	US-10-425-114-68936	Sequence 68936, A
10	1545.5	62.1	478	US-10-434-991-4	Sequence 4, Appl1
11	1545.5	62.1	478	US-10-424-599-159690	Sequence 159690, Appl1

12	1391	55.9	530	US-10-732-923-9863	Sequence 9863, Ap
13	1384.5	55.6	534	US-10-732-923-9855	Sequence 9855, Ap
14	1376.5	55.3	519	US-10-424-599-169925	Sequence 169925, Ap
15	1371	55.1	520	US-10-732-923-9856	Sequence 9856, Ap
16	1367	54.9	519	US-10-732-923-9875	Sequence 9875, Ap
17	1365.5	54.9	534	US-10-381-870-9	Sequence 9, Appl1
18	1365.5	54.9	534	US-10-732-923-9859	Sequence 9859, Ap
19	1364	54.8	532	US-10-732-923-9862	Sequence 9862, Ap
20	1347	54.1	508	US-10-732-923-9862	Sequence 9862, Ap
21	1347	54.1	518	US-10-732-923-9853	Sequence 9853, Appl1
22	1333	53.6	537	US-10-732-923-9853	Sequence 9853, Appl1
23	1325	53.3	478	US-10-437-963-156584	Sequence 156584, Ap
24	1325	53.3	478	US-10-732-923-9857	Sequence 9857, Ap
25	1325	53.3	478	US-10-732-923-9856	Sequence 9856, Ap
26	1320	53.1	536	US-10-732-923-9856	Sequence 9856, Ap
27	1316.5	52.9	490	US-10-732-923-9866	Sequence 9866, Ap
28	1316	52.9	478	US-10-381-870-4	Sequence 4, Appl1
29	1301	52.3	510	US-10-732-923-9860	Sequence 9860, Ap
30	1299.5	52.2	487	US-10-732-923-9860	Sequence 9860, Ap
31	1299.5	52.2	487	US-10-732-923-9851	Sequence 9851, Ap
32	1295.5	52.0	482	US-10-310-154-518	Sequence 518, App
33	1288	51.8	510	US-10-732-923-9867	Sequence 9867, App
34	1285.5	51.7	481	US-10-732-923-9867	Sequence 9867, App
35	1280	51.4	480	US-10-732-923-9854	Sequence 9854, App
36	1268.5	51.0	517	US-10-732-923-9854	Sequence 9854, App
37	1268.5	51.0	478	US-10-732-923-9861	Sequence 9861, App
38	1264	50.8	488	US-10-732-923-9870	Sequence 9870, App
39	1249	50.2	478	US-10-381-870-2	Sequence 2, Appl1
40	1240	49.8	512	US-10-437-963-150856	Sequence 150856, Ap
41	1193.5	48.0	511	US-10-732-923-9868	Sequence 9868, Ap
42	1193	48.0	511	US-10-732-923-9871	Sequence 9871, Ap
43	1191	47.9	488	US-10-732-923-9871	Sequence 9871, Ap
44	1186	47.7	488	US-10-732-923-9871	Sequence 9872, Ap
45	1163	46.7	514	US-10-732-923-10753	Sequence 10753, A

ALIGNMENTS

RESULT 1
US-09-884-260A-7
Sequence 7, Appl1 Application US/09884260A
Patent No. US20020098570A1
GENERAL INFORMATION:
APPLICANT: Alan Bresh
TITLE OF INVENTION: CUCUMIS MELO HYDROPEROXIDE
TITLE OF INVENTION: CUCUMIS MELO HYDROPEROXIDE
FILE REFERENCE: 06027.000202
CURRENT APPLICATION NUMBER: US/09/884.260A
PRIOR FILING DATE: 2001-06-19
PRIOR APPLICATION NUMBER: 09/537.357
NUMBER OF SEQ ID NOS: 56
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 481
TIPE: PRT
ORGANISM: Cucumis melo
US-09-884-260A-7

Query Match 100.0%; Score 2488; DB 9; Length 481;
Best Local Similarity 100.0%; Pred. No. 1.1e-233;
Matches 481; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
OY 1 MATPSSSSPPLPLKIPGCTGYGPPGAPICAKRDYFFQGRDERFRKTKNSVTRANN 60
DB 1 MATPSSSSPPLPLKIPGCTGYGPPGAPICAKRDYFFQGRDERFRKTKNSVTRANN 60
OY 61 PGPPTSSGSGPVYVLDLSPGTPPTGAPICAKRDYFFQGRDERFRKTKNSVTRANN 120
DB 61 PGPPTSSGSGPVYVLDLSPGTPPTGAPICAKRDYFFQGRDERFRKTKNSVTRANN 120

Qy 121 TEHSVVKQLFLSFLASRHRDRIPLFRSSLSSEMPVLEEDLSEKKKIADFNISISDSNSPDY 180
Db 121 TEHSVVKQLFLSFLASRHRDRIPLFRSSLSSEMPVLEEDLSEKKKIADFNISISDSNSPDY 180
Qy 181 VERLLSDGTGDSKLAEGFQMFDMLVFQIAPLASIGLPKIFSVPEDLVHTTIPLEPPPV 240
Db 181 VERLLSDGTGDSKLAEGFQMFDMLVFQIAPLASIGLPKIFSVPEDLVHTTIPLEPPPV 240
Qy 241 KSGYRLYEAFYSSSGSPIDBAEKQIDIREKCNILVFLAGFNAGCKMLPFTLLKMG 300
Db 241 KSGYRLYEAFYSSSGSPIDBAEKQIDIREKCNILVFLAGFNAGCKMLPFTLLKMG 300
Qy 301 TAGEDLHRKLAERVTIVKEEGGLTFSALRKSMLKSGVYALRIEPPVPOYGAKEDI 360
Db 301 TAGEDLHRKLAERVTIVKEEGGLTFSALRKSMLKSGVYALRIEPPVPOYGAKEDI 360
Qy 361 VIQSHDSFKIKKGETTIFGYOPATDPIFKOSKRVGDRVGESEKLLKYVMSNR 420
Db 361 VIQSHDSFKIKKGETTIFGYOPATDPIFKOSKRVGDRVGESEKLLKYVMSNR 420
Qy 421 ETVEPTANRCKCGKQLVILGRIMVPEPLRDTFTVEADLPLGPAVFKSLTRATDM 480
Db 421 ETVEPTANRCKCGKQLVILGRIMVPEPLRDTFTVEADLPLGPAVFKSLTRATDM 480
Qy 481 V 481
Db 481 V 481

RESULT 2
US-10-434-991-9
Sequence 9 Application US/10434991
Publication No US20040010822A1

GENERAL INFORMATION:
APPLICANT: McGonigle, Brian
TITLE OF INVENTION: HYDROPEROXIDE LYASES
FILE REFERENCE: BRL534USNA
CURRENT APPLICATION NUMBER: US/10/434,991
CURRENT FILING DATE: 2003-05-09
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Microsoft Word Version 7.0A
SD LENGTH: 481
TYPE: PRT
ORGANISM: Cucumis melo
US-10-434-991-9

Query Match 99.8%; Score 2463; DB 15; Length 481;
Best Local Similarity 99.8%; Pred. No. 3,46-233;

Matches 480; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MATSSSSPELPLKPIPGYGFPLGPIKIDYDYFYQGRDPEFRSRIITKNSVFRANM 60
Db 1 MATSSSSPELPLKPIPGYGFPLGPIKIDYDYFYQGRDPEFRSRIITKNSVFRANM 60
Qy 61 PRGPFISDSRNVVLDLSPILLPDTAKVERKNIIDGTWPSLSFTGNIITCAVDPSE 120
Db 61 PRGPFISDSRNVVLDLSPILLPDTAKVERKNIIDGTWPSLSFTGNIITCAVDPSE 120
Qy 121 TEHSVVKQLFLSFLASRHRDRIPLFRSSLSSEMPVLEEDLSEKKKIADFNISISDSNSPDY 180
Db 121 TEHSVVKQLFLSFLASRHRDRIPLFRSSLSSEMPVLEEDLSEKKKIADFNISISDSNSPDY 180
Qy 181 VERLLSDGTGDSKLAEGFQMFDMLVFQIAPLASIGLPKIFSVPEDLVHTTIPLEPPPV 240
Db 181 VERLLSDGTGDSKLAEGFQMFDMLVFQIAPLASIGLPKIFSVPEDLVHTTIPLEPPPV 240
Qy 241 KSGYRLYEAFYSSSGSPIDBAEKQIDIREKCNILVFLAGFNAGCKMLPFTLLKMG 300
Db 241 KSGYRLYEAFYSSSGSPIDBAEKQIDIREKCNILVFLAGFNAGCKMLPFTLLKMG 300
Qy 301 TAGEDLHRKLAERVTIVKEEGGLTFSALRKSMLKSGVYALRIEPPVPOYGAKEDI 360
Db 301 TAGEDLHRKLAERVTIVKEEGGLTFSALRKSMLKSGVYALRIEPPVPOYGAKEDI 360

Db 301 TAGEDLHRKLAERVTIVKEEGGLTFSALRKSMLKSGVYALRIEPPVPOYGAKEDI 360
Qy 361 VIQSHDSFKIKKGETTIFGYOPATDPIFKOSKRVGDRVGESEKLLKYVMSNR 420
Db 361 VIQSHDSFKIKKGETTIFGYOPATDPIFKOSKRVGDRVGESEKLLKYVMSNR 420
Qy 421 ETVEPTANRCKCGKQLVILGRIMVPEPLRDTFTVEADLPLGPAVFKSLTRATDM 480
Db 421 ETVEPTANRCKCGKQLVILGRIMVPEPLRDTFTVEADLPLGPAVFKSLTRATDM 480
Qy 481 V 481
Db 481 V 481

RESULT 3
US-09-884-260A-15
Sequence 15 Application US/09884260A
Patent No. US20020098570A1

GENERAL INFORMATION:
APPLICANT: Nathaniel Tillet
TITLE OF INVENTION: MUSCLEIN (CUCUMIS MELO) HYDROPEROXIDE
FILE REFERENCE: 06027.0002U2
CURRENT APPLICATION NUMBER: US/09/884,260A
CURRENT FILING DATE: 2001-06-19
PRIOR APPLICATION NUMBER: 09/537,357
NUMBER OF SEQ ID NOS: 56
SOFTWARE: MacSeq for Windows Version 4.0
SD LENGTH: 487
TYPE: PRT
ORGANISM: Cucumis melo
FEATURE:
NAME/KEY: VARIANT
LOCATION: (1)...(487)
OTHER INFORMATION: Xaa = Any Amino Acid
NAME/KEY: misc.feature
LOCATION: (1)...(10)
OTHER INFORMATION: Accession No. US20020098570A1 AF081955
US-09-884-260A-15

Query Match 99.0%; Score 2463.5; DB 9; Length 487;
Best Local Similarity 99.2%; Pred. No. 2,86-231;

Matches 478; Conservative 1; Mismatches 2; Indels 1; Gaps 1;

Qy 1 MATSSSSPELPLKPIPGYGFPLGPIKIDYDYFYQGRDPEFRSRIITKNSVFRANM 59
Db 1 MATSSSSPELPLKPIPGYGFPLGPIKIDYDYFYQGRDPEFRSRIITKNSVFRANM 60
Qy 60 MPGPFISDSRNVVLDLSPILLPDTAKVERKNIIDGTWPSLSFTGNIITCAVDPSE 119
Db 60 MPGPFISDSRNVVLDLSPILLPDTAKVERKNIIDGTWPSLSFTGNIITCAVDPSE 120
Qy 120 TEHSVVKQLFLSFLASRHRDRIPLFRSSLSSEMPVLEEDLSEKKKIADFNISISDSNSPDY 179
Db 120 TEHSVVKQLFLSFLASRHRDRIPLFRSSLSSEMPVLEEDLSEKKKIADFNISISDSNSPDY 180
Qy 180 YVPELLSDGTGDSKLAEGFQMFDMLVFQIAPLASIGLPKIFSVPEDLVHTTIPLEPPPV 239
Db 180 YVPELLSDGTGDSKLAEGFQMFDMLVFQIAPLASIGLPKIFSVPEDLVHTTIPLEPPPV 240
Qy 240 YVSGYRLYEAFYSSSGSPIDBAEKQIDIREKCNILVFLAGFNAGCKMLPFTLLKMG 299
Db 240 YVSGYRLYEAFYSSSGSPIDBAEKQIDIREKCNILVFLAGFNAGCKMLPFTLLKMG 300
Qy 300 GRGPEDLHRKLAERVTIVKEEGGLTFSALRKSMLKSGVYALRIEPPVPOYGAKEDI 359
Db 301 GRGPEDLHRKLAERVTIVKEEGGLTFSALRKSMLKSGVYALRIEPPVPOYGAKEDI 360
Qy 360 YIQSHDSFKIKKGETTIFGYOPATDPIFKOSKRVGDRVGESEKLLKYVMSNR 419
Db 360 YIQSHDSFKIKKGETTIFGYOPATDPIFKOSKRVGDRVGESEKLLKYVMSNR 419

Db 361 IVQSHSSFKIKGGTIGQPPATNDPKIFDSEKFDGDRPVSGSEKLTATVWSN 420
Qy 420 RETVEPTAKKQCGKRLVYLGRIWVEFFLARDFTFVADLPLGPAVKKSLTRATD 479
Db 421 RETVEPTAKKQCGKRLVYLGRIWVEFFLARDFTFVADLPLGPAVKKSLTRATD 480
Qy 480 MV 481
Db 481 ML 482

RESULT 4
US-10-434-991-10
Sequence 10, Application US/10434991
Publication No. US20040010822A1
Applicant: Mcgonigle, Brian
Title of Invention: HYDROPEROXIDE LYASES
File Reference: BB1534USNA
Current Application Number: US/10/434,991
Current Filing Date: 2003-05-09
Number of Seq ID Nos: 14
Software: Microsoft Word Version 7.0A
Seq ID No 10
Length: 478
Type: PRT
Organism: Cucumis sativus
US-10-434-991-10

Query Match 94.0%; Score 2339; DB 15; Length 478;
Best Local Similarity 93.9%; Pred. No. 3,90-219;
Matches 448; Conservative 14; Mismatches 15; Indels 0; Gaps 0;

Qy 5 SSSSPELPKPIPGGYFPPLGPIKORVDYFYQGRDSEFRSRTTKYNSIVFRANMPGP 64
Db 2 ASSSPELPKPIPGGYFPPLGPIKORVDYFYQGRDSEFRSRTTKYNSIVFRANMPGP 61
Qy 65 FISDSRVVYLLDALSPFLLPDTAKVERKNTLDGTYNPSLSFTGNITFCAYLDSESTHS 124
Db 62 FISDSRVVYLLDALSPFLLPDTAKVERKNTLDGTYNPSLSFTGNITFCAYLDSESTHS 121
Qy 125 VAKRLPFLASRHDHFPIPLFRSSLSBMPKVEDLSKKKIDAPNSISDSMSFDYVRL 184
Db 122 VAKRLPFLASRHDHFPIPLFRSSLSBMPKVEDLSKKKIDAPNSISDSMSFDYVRL 181
Qy 185 LSDGTFDKLAAGKGMFDLMVFOPLASIGLKIIFVPEDLVHTIPLPPPPVSGY 244
Db 182 LSDGTFDKLAAGKGMFDLMVFOPLASIGLKIIFVPEDLVHTIPLPPPPVSGY 241
Qy 245 RLKYEAFYSSSGSFLDEAKQIDREKACNVLVFLAGNAYGKVLFPFLKRWGTAGE 304
Db 242 RLKYEAFYSSSGSFLDEAKQIDREKACNVLVFLAGNAYGKVLFPFLKRWGTAGE 301
Qy 305 DLHAKLAERTVTVKGGCTFSALTKRSLKSYVYVYALRTEPPVPOYGAKEDIV 364
Db 302 DLHAKLAERTVTVKGGCTFSALTKRSLKSYVYVYALRTEPPVPOYGAKEDIV 361
Qy 365 HDSEFKIKKGGTIFGYOPFATDPKIFDSEKFDGDRPVSGSEKLTATVWSN 424
Db 362 HDSEFKIKKGGTIFGYOPFATDPKIFDSEKFDGDRPVSGSEKLTATVWSN 421
Qy 425 PTANRQCGKGNLVYLGRIMVVEFFLARDFTFVADLPLGPAVKKSLTRATDV 481
Db 422 PTANRQCGKGNLVYLGRIMVVEFFLARDFTFVADLPLGPAVKKSLTRATDV 478

RESULT 5
US-10-686-947-156
Sequence 156, Application US/10686947
Publication No. US20040162420A1
Applicant: Profigen Inc.

FILE REFERENCE: 79601
CURRENT APPLICATION NUMBER: US/10/686,947
CURRENT FILING DATE: 2003-10-16
PRIOR APPLICATION NUMBER: US/10/387346
PRIOR FILING DATE: 2003-03-12
NUMBER OF SEQ ID NOS: 298
SOFTWARE: PatentIn version 3.2
Seq ID No 156
Length: 483
Type: PRT
Organism: NICOTIANA GLABRA
US-10-686-947-156

Query Match 64.9%; Score 1613.5; DB 16; Length 483;
Best Local Similarity 63.9%; Pred. No. 2,90-148;
Matches 305; Conservative 75; Mismatches 92; Indels 5; Gaps 4;

Qy 5 SSSSPELPKPIPGGYFPPLGPIKORVDYFYQGRDSEFRSRTTKYNSIVFRANMPGP 64
Db 9 ASSSPELPKPIPGGYFPPLGPIKORVDYFYQGRDSEFRSRTTKYNSIVFRANMPGP 68
Qy 65 FISDSRVVYLLDALSPFLLPDTAKVERKNTLDGTYNPSLSFTGNITFCAYLDSESTHS 124
Db 62 FISDSRVVYLLDALSPFLLPDTAKVERKNTLDGTYNPSLSFTGNITFCAYLDSESTHS 128
Qy 125 VAKRLPFLASRHDHFPIPLFRSSLSBMPKVEDLSKKKIDAPNSISDSMSFDYVRL 184
Db 122 VAKRLPFLASRHDHFPIPLFRSSLSBMPKVEDLSKKKIDAPNSISDSMSFDYVRL 187
Qy 185 LSDGTFDKLAAGKGMFDLMVFOPLASIGLKIIFVPEDLVHTIPLPPPPVSGY 244
Db 182 LSDGTFDKLAAGKGMFDLMVFOPLASIGLKIIFVPEDLVHTIPLPPPPVSGY 247
Qy 245 RLKYEAFYSSSGSFLDEAKQIDREKACNVLVFLAGNAYGKVLFPFLKRWGTAGE 304
Db 242 RLKYEAFYSSSGSFLDEAKQIDREKACNVLVFLAGNAYGKVLFPFLKRWGTAGE 307
Qy 305 DLHAKLAERTVTVKGGCTFSALTKRSLKSYVYVYALRTEPPVPOYGAKEDIV 364
Db 302 DLHAKLAERTVTVKGGCTFSALTKRSLKSYVYVYALRTEPPVPOYGAKEDIV 367
Qy 365 HDSEFKIKKGGTIFGYOPFATDPKIFDSEKFDGDRPVSGSEKLTATVWSN 424
Db 362 HDSEFKIKKGGTIFGYOPFATDPKIFDSEKFDGDRPVSGSEKLTATVWSN 426
Qy 425 PTANRQCGKGNLVYLGRIMVVEFFLARDFTFVADLPLGPAVKKSLTRATDV 481
Db 422 PTANRQCGKGNLVYLGRIMVVEFFLARDFTFVADLPLGPAVKKSLTRATDV 478
Qy 427 TDPVFNKQCGKGNLVYLGRIMVVEFFLARDFTFVADLPLGPAVKKSLTRATDV 483

RESULT 6
US-10-434-991-5
Sequence 5, Application US/10434991
Publication No. US20040010822A1
Applicant: Mcgonigle, Brian
Title of Invention: HYDROPEROXIDE LYASES
File Reference: BB1534USNA
Current Application Number: US/10/434,991
Current Filing Date: 2003-05-09
Number of Seq ID Nos: 14
Software: Microsoft Word Version 7.0A
Seq ID No 6
Type: PRT
Organism: Glycine max
US-10-434-991-5

Query Match 62.7%; Score 1560; DB 15; Length 487;
Best Local Similarity 60.3%; Pred. No. 50-143;
Matches 291; Conservative 76; Mismatches 107; Indels 4; Gaps 3;

QY 183 RLISDGT PDSK--LAABGPGMFDLWLVQLAPLASIGLPKIFSVFEDLVHTIPLPPPV 240

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Db      224  RLCGCKDPSQTNIGSSGKLVDPVNLPLQALPAPLALCEKRIINPIIDPLIPPLPPACL 283
Oy      241  KSGYKRYLVEAFVSSGSLPLDEAKKQGIIDERKACNLTVPAGFNAYGKMKLPFTLLKRWG 300
Db      284  KSSYKYNLEAFVFTATYALDBAKGLKLRKQACNHNVPFAGNAYGGLKNDPPYALKWLD 343
Oy      301  TAGEQDLRKALAPRTTVEYSGLTSTAKKQSLIKSVYVKAALIRPPPIPGYAKAZDI 360
Db      344  LSGEKLADLAIEARVAVDDSGVFTLLENPVLTKSVYEVYEMHISPAVPQYADARNL 403
Oy      361  VIGSHDSFPIKKEGTIPQYPPATADKLPKXDSKKVQDRVVEEGELKATKVMNSR 420
Db      404  VVSHSDSPFVYKQKMLPGQYPPATSDPRIPEDAEVAVRRVRPG-GEKMLKIVLMSNG 462
Oy      421  ETVEPFAKRCQPSQNLVYLIGRIWVEFFLARYDFTVEVADPLGPAVKESLTPAT 478
Db      463  ETSEFSAKRCQPSQNLVVLCLRLPVELRLDIFSEFTYQAGFPTITTLIKTNS 520

RESULT 8
US-10-425-114-40766
* Sequence 40766, Application US/10425114
* Publication No. US20040034888A1
* GENERAL INFORMATION:
* APPLICANT: LAY, Jingsong
* APPLICANT: Zhou, Jihua
* APPLICANT: Zhou, David K.
* APPLICANT: Screen, Steven B
* APPLICANT: Tabaska, Jack B
* APPLICANT: Cao, Yongwei
* TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
* TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
* FILE REFERENCE: 38-21(53313) B
* CURRENT APPLICATION NUMBER: US/10-425-114
* CURRENT FILING DATE: 2003-04-28
* NUMBER OF SEQ ID NOS: 73128
* SEQ ID NOS: 40766
* TYPE: PRT
* LENGTH: 492
* ORGANISM: Glycine max
* FEATURE:
* OTHER INFORMATION: Clone ID: 7007323423_F11.pep
* US-10-425-114-40766

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[illegible]

[illegible][illegible]

Query Match	62.1%	Score 1545.5;	DB 15;	Length 478;
Best Local Similarity	59.8%	Pred. No. 1.3e-14;		
Matches 284; Conservative	88;	Mismatches 102;	Indels 1;	Gaps 1;

[illegible]

3 TPSSSP---ELPLKIPGVGPPPLDIPKQKDYVYFGQDRFFSRRTITNNTVYPA 58
 :::
 Overall Match Score 55.9%; Score 1391; DB-17; Length 530;
 Best Local Similarity 55.8%; Pred. No. 1:8-126;
 Matches 269; Conservative 80; Mismatches 123; Indels 10; Gaps 5

Dh	52	SPFSSPVPKQAKLPKRVKGVQGLVLPKMPKDBLDFPNQGNKRFPSKSIQKQSTVART	111
Qy	59	NMPGGPFTSSDSQSVVYLVLDALSPILLTPPAKVEKRNVLIDTTPBLSPTGNITPCYALAD	116
Dh	112	NMPGGPFTSPRVVYLVLDALSPILLTPPAKVEKRNVLIDTTPBLSPTGNITPCYALAD	117
Qy	119	SPFTSPVYKGLFSLASHSHRRTFPLSPKSPKQGLTGKSTREKXIDLVSGISVSLD	176
Dh	172	SPSPHAKKMLPVYLSPPRRVYIEFHNSVSELFTEDENELSTCKG-ARLNAANDQAF	230
Qy	179	DYVRYLLIDDTG-DSCKAALBPQMDLMLVPLAPLASIGLPIYVSFVSDVLIHTPIAP	237
Dh	231	NFLANLBYGINODPTKGLDQPLGKMWLPOLHPIILLIGLPG--VLEDDLVNHTRLAP	287
Qy	238	PVPSGYKRLYEAFPSSSGFLDLEAKQIGIDREKACMLVFLANGFVAYSKVLPLILK	297
Dh	286	ALVYKQDQLRLNFPKNSVLYDLEKELTGISSEKSCMLHPTATCSFGSGLTIFPANKL	344
Qy	298	NMGTAGDDELRKLEKRYFTWYEEEG-LTPSKALEKRYSLASVYELALIPBPQYQXGA	355
Dh	348	WIGRAQALISQMDQETRSVYSSNGKVMYAMKQPMKSVYVESLRLIEPPVAQYGRA	407
Qy	357	KEDLVIGSHSSPKKKGGTIFGYQFPATADPKI FDSGEKFGQDFPVGSEGEELKLVYV	416
Dh	408	KIDVYIEHSHASBELKGGVLYGFQFPATADPKI FDSGEKFGQDFPVGSEGEELKLVYV	467
Qy	417	SNEETVEFPANNOCKGNLVYLIGRIWVEFPLKDYITPVEVADLPLGAVKRFSLTR	476
Dh	476	SNGETVEFNFSINNOCKGNQDPPVYVSKLIDVPLKDYITPVEVADLPLGAVKRFSLTR	527
Qy	477	AS	529
Dh	528	AS	529
RESULT 13			
US-10-732-923-9855			
Sequence 9855, Application US/107329923			
Publ. No. US/20050108791X1			
Publ. Date 11/11/2005			
Applicant: Edgerton, Michael D			
Title of Invention: TRANSGENIC PLANTS WITH IMPROVED PHENOTYPES			
File Reference: 38-15(527961C)			
Current Application Number: US/10/732,923			
Current Filing Date: 2003-12-10			
Prior Application Number: 10/310,154			

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? SEQ ID NO 9855
? LENGTH: 534
? TYPE: PRT Glycine max
? OS: Glycine max
? US-10-732-923-9855

Query Match      55.6%; Score 1384.5; DB 17; Length 534;
Best Local Similarity 54.7%; Pred. NO. 98-126;
Matches 262; Conservative 86; Mismatches 122; Indels 9; Gaps 5;

OY 5 SSSSRF...LAKPIFGQYQGLPGLPIKRDYDFIRQGRDEFKRSRTKNSYVRAMP 61
Db 59 STSREKSLIKIKIGNOCPVIGDKKDODIFYQGRDEFKRSIKQKSTVPRITM 118

OY 62 PGPFISDSRVRVYLALASPLILPRTAKRKRNILDGYWPSLSTNIRFCALDPER 121
Db 119 PGFGLANRRVYVYLADATAPLILPGRKSKVDGVRGPGSTQITQGYRLASLDPER 178

OY 122 EHSYTKLFLSLASHDRRIPLFRSLSEMPVVDLSEKKIKLADPNISLSDNSP 181
Db 179 KISIKLQFLFLSKRRAPVISEPHASVYDQFHALDNLAEKKLASFQDNOAAVPL 230

OY 182 FRLSLDQTP-DSKIAAGPGQFDMLVQLPVLASLGLPRTFSYEDLVYITPLPP 247
Db 238 SRELNSNPNADTKRGSDPKIKVQKMLFQLPILRUSGDP--LSESTISRLPSL 294

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Oy 241 KGTGKLYEAFTYSSGSLDPAKQIDREKACINLVPLAGFNAYGCKKLPFTLLKRW 300
Db 295 QKDTQMLDFTFFQSSGVYDDBAKLQITNDBKCHMLVTCFMSFGCKMLPFPVLAHIG 354
Oy 301 TAGEDLHRLKLAERTVTVKESG-LTFSALEKMSLKSVYVBLRTEPPVPQYGAKE 359
Db 355 RAGVYKLAHRLAEIRSAVAGCEITMAAEMMPAKSVYVEAFRIDPEVPYQFGAKED 414
Oy 360 IVTQSHDSFKIKKGETIPGYOPATKDPKIFKQSEKVFQDRFVGESEKLLKTVYWSN 419
Db 415 LIIESHDAHFOVEGEMLPGYOPATKDPKIFERAEFVQDRFVGESEKLLKTVYWSN 474
Oy 420 RETVEFPAENKQCGKMLVILGRIMVVEFPLRDTFTVEVADLPLGPAVFKSLTRAT 478
Db 475 PBTSEFTIGNKQAGKDPVTLVSRLLVVEFPLRDTFTVEVADLPLGPAVFKSLTRAT 533

RESULT 14
US-10-424-599-169925
Sequence 169925, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
FILE REFERENCE: 38-21(4322) B
CURRENT APPLICATION NUMBER: US/10/424,599
PRIOR FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 169925
LENGTH: 519
TYPE: PRT
ORGANISM: Glycine max
FEATURE:
OTHER INFORMATION: Clone ID: PAT MRT3841_124457C.1.pcp
US-10-424-599-169925

Query Match 55.3%; Score 1376.5; DB 15; Length 519;
Best Local Similarity 54.5%; Pred. No. 4,66-125;
Matches 261; Conservative 85; Mismatches 124; Indels 9; Gaps 5;

Oy 5 SSSPPE--LPKPIPGGYGPPPLGPIKORADYFPGQDEFPKRSITKTSVFRANP 61
Db 44 SVTSPPEKLPKIRKIPDGGFVYGLDRODPIFKQGBDFPKSKIOKQSVFRANP 103
Oy 62 RGPPISSDSRV-VYLDALSPILPDTAKTEKNLIDGTWBSLSTGNIRTCAYLDPS 121
Db 104 RGPPLADPNVYVYLDLAKSPFLPDSNKKYKQVDTGTMTSTLTCGYVSLYLDPS 163
Oy 122 HSTVLAHRLFLSLAERHDPFLPLFRSLSEMPVKLEKSEKKIADPNISIDPSFDY 181
Db 164 KHALAQMLPFLKSRBAVYISFHASYKELPHALENLAENK-ASFGDANDQAAENF 222
Oy 182 PALLSGDTP-DSKLAAGPQMDLWVQLAPLAGIGLPIFSVFEVADLPLGPAVFKSLTRAT 240
Db 223 SRSLSFNSNADTLGIDGPRIVQMWLPQIGPLMLDLPQ--LDSSTIRFLPSL 279
Oy 241 KGTGKLYEAFTYSSGSLDPAKQIDREKACINLVPLAGFNAYGCKKLPFTLLKRW 300
Db 280 QKDTQMLDFTFFQSSGVYDDBAKLQITNDBKCHMLVTCFMSFGCKMLPFPVLAHIG 354
Oy 301 TAGEDLHRLKLAERTVTVKESG-LTFSALEKMSLKSVYVBLRTEPPVPQYGAKE 359
Db 340 RAGVYKLAHRLAEIRSAVAGCEITMAAEMMPAKSVYVEAFRIDPEVPYQFGAKED 414
Oy 360 IVTQSHDSFKIKKGETIPGYOPATKDPKIFKQSEKVFQDRFVGESEKLLKTVYWSN 419
Db 415 LIIESHDAHFOVEGEMLPGYOPATKDPKIFERAEFVQDRFVGESEKLLKTVYWSN 474

Oy 420 RETVEFPAENKQCGKMLVILGRIMVVEFPLRDTFTVEVADLPLGPAVFKSLTRAT 478
Db 460 PBTSEFTIGNKQAGKDPVTLVSRLLVVEFPLRDTFTVEVADLPLGPAVFKSLTRAT 533

RESULT 15
US-10-732-923-9856
Sequence 9856, Application US/10732923
Publication No. US20050108791A1
GENERAL INFORMATION:
APPLICANT: Edgerton, Michael D
TITLE OF INVENTION: TRANSLOC PLANTS WITH IMPROVED PHENOTYPES
FILE REFERENCE: 38-21(4322) B
CURRENT APPLICATION NUMBER: US/10/732,923
PRIOR FILING DATE: 2003-12-10
PRIOR FILING DATE: 2002-12-04
NUMBER OF SEQ ID NOS: 24149
SEQ ID NO 9856
LENGTH: 520
TYPE: PRT
ORGANISM: Glycine max
FEATURE:
NAME/KEY: unsure
LOCATOR: 11...1620
OTHER INFORMATION: unsure at all Kna locations
US-10-732-923-9856

Query Match 55.1%; Score 1371; DB 17; Length 520;
Best Local Similarity 54.6%; Pred. No. 1,66-124;
Matches 262; Conservative 83; Mismatches 125; Indels 10; Gaps 6;

Oy 5 SSSPPE--LPKPIPGGYGPPPLGPIKORADYFPGQDEFPKRSITKTSVFRANP 61
Db 44 SVTSPPEKLPKIRKIPDGGFVYGLDRODPIFKQGBDFPKSKIOKQSVFRANP 103
Oy 62 RGPPISSDSRV-VYLDALSPILPDTAKTEKNLIDGTWBSLSTGNIRTCAYLDPS 121
Db 104 RGPPLADPNVYVYLDLAKSPFLPDSNKKYKQVDTGTMTSTLTCGYVSLYLDPS 163
Oy 122 HSTVLAHRLFLSLAERHDPFLPLFRSLSEMPVKLEKSEKKIADPNISIDPSFDY 181
Db 164 KHALAQMLPFLKSRBAVYISFHASYKELPHALENLAENK-ASFGDANDQAAENF 222
Oy 182 PALLSGDTP-DSKLAAGPQMDLWVQLAPLAGIGLPIFSVFEVADLPLGPAVFKSLTRAT 240
Db 223 SRSLSFNSNADTLGIDGPRIVQMWLPQIGPLMLDLPQ--LDSSTIRFLPSL 279
Oy 241 KGTGKLYEAFTYSSGSLDPAKQIDREKACINLVPLAGFNAYGCKKLPFTLLKRW 300
Db 280 QKDTQMLDFTFFQSSGVYDDBAKLQITNDBKCHMLVTCFMSFGCKMLPFPVLAHIG 354
Oy 301 TAGEDLHRLKLAERTVTVKESG-LTFSALEKMSLKSVYVBLRTEPPVPQYGAKE 359
Db 340 RAGVYKLAHRLAEIRSAVAGCEITMAAEMMPAKSVYVEAFRIDPEVPYQFGAKED 414
Oy 360 IVTQSHDSFKIKKGETIPGYOPATKDPKIFKQSEKVFQDRFVGESEKLLKTVYWSN 419
Db 415 LIIESHDAHFOVEGEMLPGYOPATKDPKIFERAEFVQDRFVGESEKLLKTVYWSN 474

Search completed: October 5, 2005, 07:13:01
Job time : 170 secs